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OM protein - protein search, using sw model

Run on: May 19, 2003, 16:43:42 ; Search time 19.8719 Seconds  
(without alignments)  
553.754 Million cell updates/sec

Title: US-09-625-573-2  
Perfect score: 1970  
Sequence: 1 MLSTSRFRIRNTNESGEV.....GKGKSIGRAPEASLQDKEGA 374

Scoring table: BLOSUM62  
Gapop 10.0 , Gapext 0.5

Searched: 262574 seqs, 29422922 residues  
Total number of hits satisfying chosen parameters: 262574

Minimum DB seq length: 0  
Maximum DB seq length: 2000000000  
Post-processing: Minimum Match 0%  
Maximum Match 100%  
Listing first 45 summaries

Database : Issued Patents, AA: \*  
1: /cgn2\_6/ptodata/1/iaa/5A\_COMB.pep.\*  
2: /cgn2\_6/ptodata/1/iaa/5B\_COMB.pep.\*  
3: /cgn2\_6/ptodata/1/iaa/6A\_COMB.pep.\*  
4: /cgn2\_6/ptodata/1/iaa/6B\_COMB.pep.\*  
5: /cgn2\_6/ptodata/1/iaa/PTUS\_COMB.pep.\*  
6: /cgn2\_6/ptodata/1/iaa/backfiles1.pep.\*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

## SUMMARIES

Result No.	Score	Query Match %	Length	DB ID	Description
1	1970	100.0	374	1	US-08-450-393A-2
2	1970	100.0	374	4	US-08-446-669-2
3	1970	100.0	374	5	PCT-US95-00476-2
4	1823	92.5	344	3	US-08-466-343D-9
5	1651.5	83.8	360	1	US-08-450-393A-4
6	1651.5	83.8	360	4	US-08-446-669-4
7	1651.5	83.8	360	4	US-09-045-583-50
8	1651.5	83.8	360	4	US-09-534-185-50
9	1645.5	83.5	360	5	PCT-US95-00476-4
10	1645.5	83.5	360	4	US-08-833-752-7
11	1614.5	82.0	360	4	US-09-045-583-51
12	1614.5	82.0	360	4	US-09-534-185-51
13	1589.5	80.7	347	1	US-08-461-244-3
14	1236	62.7	352	4	US-09-517-605-5
15	1234	62.6	354	4	US-08-724-984A-2
16	1230	62.4	352	4	US-09-045-583-52
17	1224	62.4	352	4	US-09-534-185-52
18	1224	62.1	352	4	US-09-087-232A-13
19	1224	62.1	352	4	US-08-861-105-14
20	1224	62.1	352	4	US-08-575-967A-2
21	1224	62.1	352	4	US-08-833-752-5
22	1215	61.7	352	3	US-08-466-343D-2
23	967.5	49.1	355	1	US-08-012-988A-2
24	967.5	49.1	355	1	US-08-450-393A-5
25	967.5	49.1	355	4	US-08-446-669-5
26	967.5	49.1	355	4	US-09-239-938-1
27	967.5	49.1	355	5	PCT-US95-00476-5

Sequence 9, Appli  
Sequence 53, Appl  
Sequence 53, Appl  
Sequence 4, Appli  
Sequence 1, Appli  
Sequence 54, Appl  
Sequence 8, Appli  
Sequence 20, Appl  
Sequence 2, Appli  
Sequence 55, Appl  
Sequence 55, Appl  
Sequence 10, Appl  
Sequence 2, Appli  
Sequence 56, Appl  
Sequence 56, Appl  
Sequence 28, Appl  
Sequence 32, Appl

28 940.5 47.7 355 4 US-08-833-752-9  
29 911.5 46.3 355 4 US-09-045-583-53  
30 911.5 46.3 355 4 US-09-534-185-53  
31 886.5 45.0 355 4 US-08-575-967A-4  
32 886.5 45.0 355 4 US-08-847-296B-1  
33 886.5 45.0 355 4 US-09-045-583-54  
34 886.5 45.0 355 4 US-09-534-185-54  
35 858.5 43.6 355 4 US-08-833-752-8  
36 831.5 42.2 360 4 US-08-875-573-20  
37 831.5 42.2 360 4 US-09-232-878-2  
38 831.5 42.2 360 4 US-09-045-583-55  
39 831.5 42.2 360 4 US-09-534-185-55  
40 797 40.5 360 4 US-08-833-752-10  
41 723 36.7 355 1 US-08-461-244-2  
42 723 36.7 355 4 US-09-045-583-56  
43 723 36.7 355 4 US-09-534-185-56  
44 698 35.4 355 1 US-08-153-848-28  
45 698 35.4 355 1 US-08-153-848-32

## ALIGNMENTS

RESULT 1  
US-08-450-393A-2  
; Sequence 2, Application US/08450393A  
; Patent No. 5707815  
; GENERAL INFORMATION:  
; APPLICANT: Charo, Israel  
; APPLICANT: Coughlin, Shaun  
; TITLE OF INVENTION: MAMMALIAN MONOCYTE CHEMOATTRACTANT  
; TITLE OF INVENTION: PROTEIN RECEPTORS  
; NUMBER OF SEQUENCES: 14  
; CORRESPONDENCE ADDRESS:  
; ADDRESS: Cooley Godward Castro Huddleson & Tatum  
; STREET: 5 Palo Alto Square  
; CITY: Palo Alto  
; STATE: California  
; COUNTRY: USA  
; ZIP: 94306-2155  
; COMPUTER READABLE FORM:  
; MEDIUM TYPE: Floppy disk  
; COMPUTER: IBM PC compatible  
; OPERATING SYSTEM: PC-DOS/MS-DOS  
; SOFTWARE: Patent Release #1.0, Version #1.25  
; CURRENT APPLICATION DATA:  
; APPLICATION NUMBER: US/08/450,393A  
; FILING DATE: May 25, 1995  
; CLASSIFICATION: 424  
; ATTORNEY/AGENT INFORMATION:  
; NAME: Cserr, Luann  
; REGISTRATION NUMBER: 31,822  
; REFERENCE/DOCKET NUMBER: UCAL-237/02US  
; TELECOMMUNICATION INFORMATION:  
; TELEPHONE: 415-843-5165  
; TELEFAX: 415-8857-0663  
; TELEX: 380816CooleyPA  
; INFORMATION FOR SEQ ID NO: 2:  
; SEQUENCE CHARACTERISTICS:  
; LENGTH: 374 amino acids  
; TYPE: amino acid  
; TOPOLOGY: linear  
; MOLECULE TYPE: protein  
US-08-450-393A-2

Query Match 100.0%; Score 1970; DB 1; Length 374;  
Best Local Similarity 100.0%; Pred. No. 1.8e-150;  
Matches 374; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 MLSTSRFRIRNTNESGEVTTFFDYDYGAPCHKFDVKGQIAGQLLPPLYSLVFIFGVGN 60  
Db 1 MLSTSRFRIRNTNESGEVTTFFDYDYGAPCHKFDVKGQIAGQLLPPLYSLVFIFGVGN 60

QY 61 MLVVLILINCKKLCITDIYLLNLAIISDLLELITLPLWAHSAANEWVFGNAMCKLFTGLY 120  
DB 61 MLVVLILINCKKLCITDIYLLNLAIISDLLELITLPLWAHSAANEWVFGNAMCKLFTGLY 120  
QY 121 HIGYFGGIFFIILLTIDRYLAIVHAFKARTVTFGVVTSVITWLVAFAVSPGIIFTK 180  
DB 121 HIGYFGGIFFIILLTIDRYLAIVHAFKARTVTFGVVTSVITWLVAFAVSPGIIFTK 180  
QY 181 COKEDSVYCGPYPPRGWNNFHTIMRNILGLVPLLMVICYSGILKTLRCRNEKKRHR 240  
DB 181 COKEDSVYCGPYPPRGWNNFHTIMRNILGLVPLLMVICYSGILKTLRCRNEKKRHR 240  
QY 241 AVRVIETIMIVYFLWTPYNYVILLNTFOEFFGLSNCESTSQLDQATQVETLGMTHCCI 300  
DB 241 AVRVIETIMIVYFLWTPYNYVILLNTFOEFFGLSNCESTSQLDQATQVETLGMTHCCI 300  
QY 301 NPIIYAFVGEKFRSLFHIALGCRIPALQKPVCGGPGVPRGNKVKVTTQGLLDGRGKXSI 360  
DB 301 NPIIYAFVGEKFRSLFHIALGCRIPALQKPVCGGPGVPRGNKVKVTTQGLLDGRGKXSI 360  
361 GRAPEASLQDKEGA 374  
DB 361 GRAPEASLQDKEGA 374

## RESULT 2

US-08-446-669-2  
; Sequence 2, Application US/08446669  
; Patent No. 6132987  
; GENERAL INFORMATION:  
; APPLICANT: Charo, Israel  
; APPLICANT: Coughlin, Shaun  
; TITLE OF INVENTION: MAMMALIAN MONOCYTE CHEMOATTRACTANT  
; TITLE OF INVENTION: PROTEIN RECEPTORS  
; NUMBER OF SEQUENCES: 14  
; CORRESPONDENCE ADDRESS:  
; ADDRESSEE: Cooley Godward Castro Huddleson & Tatum  
; STREET: 5 Palo Alto Square  
; CITY: Palo Alto  
; STATE: California  
; COUNTRY: USA  
; ZIP: 94306-2155  
; COMPUTER READABLE FORM:  
; MEDIUM TYPE: Floppy disk  
; COMPUTER: IBM PC compatible  
; OPERATING SYSTEM: PC-DOS/MS-DOS  
; SOFTWARE: Patent In Release #1.0, Version #1.25  
; CURRENT APPLICATION DATA:  
; APPLICATION NUMBER: US/08/446,669  
; FILING DATE: May 25, 1995  
; CLASSIFICATION: 435  
; ATTORNEY/AGENT INFORMATION:  
; NAME: Neeley, Richard  
; REGISTRATION NUMBER: 30,092  
; REFERENCE/DOCKET NUMBER: UCAL-237/01US  
; TELECOMMUNICATION INFORMATION:  
; TELEPHONE: 415-843-5000  
; TELEFAX: 415-857-0663  
; TELEX: 380816COOLEYPA  
; INFORMATION FOR SEQ ID NO: 2:  
; SEQUENCE CHARACTERISTICS:  
; LENGTH: 374 amino acids  
; TYPE: amino acid  
; TOPOLOGY: linear  
; MOLECULE TYPE: protein  
US-08-446-669-2

Query Match 100.0%; Score 1970; DB 4; Length 374;  
Best Local Similarity 100.0%; Pred. No. 1.8e-150;  
Matches 374; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 MLSTSRSRFIRNTNESGEVTTFFDYDYGAPCHKFDVKQIGAOQLLPPLYSLVFIQFVGN 60  
|||||

DB 1 MLSTSRSRFIRNTNESGEVTTFFDYDYGAPCHKFDVKQIGAOQLLPPLYSLVFIQFVGN 60  
QY 61 MLVVLILINCKKLCITDIYLLNLAIISDLLELITLPLWAHSAANEWVFGNAMCKLFTGLY 120  
DB 61 MLVVLILINCKKLCITDIYLLNLAIISDLLELITLPLWAHSAANEWVFGNAMCKLFTGLY 120  
QY 121 HIGYFGGIFFIILLTIDRYLAIVHAFKARTVTFGVVTSVITWLVAFAVSPGIIFTK 180  
DB 121 HIGYFGGIFFIILLTIDRYLAIVHAFKARTVTFGVVTSVITWLVAFAVSPGIIFTK 180  
QY 181 COKEDSVYCGPYPPRGWNNFHTIMRNILGLVPLLMVICYSGILKTLRCRNEKKRHR 240  
DB 181 COKEDSVYCGPYPPRGWNNFHTIMRNILGLVPLLMVICYSGILKTLRCRNEKKRHR 240  
QY 241 AVRVIETIMIVYFLWTPYNYVILLNTFOEFFGLSNCESTSQLDQATQVETLGMTHCCI 300  
DB 241 AVRVIETIMIVYFLWTPYNYVILLNTFOEFFGLSNCESTSQLDQATQVETLGMTHCCI 300  
QY 301 NPIIYAFVGEKFRSLFHIALGCRIPALQKPVCGGPGVPRGNKVKVTTQGLLDGRGKXSI 360  
DB 301 NPIIYAFVGEKFRSLFHIALGCRIPALQKPVCGGPGVPRGNKVKVTTQGLLDGRGKXSI 360  
361 GRAPEASLQDKEGA 374  
DB 361 GRAPEASLQDKEGA 374

## RESULT 3

PCT-US95-00476-2  
; Sequence 2, Application PC/TUS9500476  
; GENERAL INFORMATION:  
; APPLICANT: The Regents of the University of California  
; TITLE OF INVENTION: MAMMALIAN MONOCYTE CHEMOATTRACTANT  
; TITLE OF INVENTION: PROTEIN RECEPTORS  
; NUMBER OF SEQUENCES: 14  
; CORRESPONDENCE ADDRESS:  
; ADDRESSEE: Robbins, Berliner & Carson  
; STREET: 201 N. Figueroa Street, 5th Floor  
; CITY: Los Angeles  
; STATE: California  
; COUNTRY: USA  
; ZIP: 90012-2628  
; COMPUTER READABLE FORM:  
; MEDIUM TYPE: Floppy disk  
; COMPUTER: IBM PC compatible  
; OPERATING SYSTEM: PC-DOS/MS-DOS  
; SOFTWARE: Patent In Release #1.0, Version #1.25  
; CURRENT APPLICATION DATA:  
; APPLICATION NUMBER: PCT/US95/00476  
; FILING DATE:  
; CLASSIFICATION:  
; ATTORNEY/AGENT INFORMATION:  
; NAME: Berliner, Robert  
; REGISTRATION NUMBER: 20,121  
; REFERENCE/DOCKET NUMBER: 5555-291  
; TELECOMMUNICATION INFORMATION:  
; TELEPHONE: 310-977-1001  
; TELEFAX: 310-977-1003  
; TELEX:  
; INFORMATION FOR SEQ ID NO: 2:  
; SEQUENCE CHARACTERISTICS:  
; LENGTH: 374 amino acids  
; TYPE: amino acid  
; TOPOLOGY: linear  
; MOLECULE TYPE: protein  
PCT-US95-00476-2

Query Match 100.0%; Score 1970; DB 5; Length 374;  
Best Local Similarity 100.0%; Pred. No. 1.8e-150;  
Matches 374; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 MLSTSRSRFIRNTNESGEVTTFFDYDYGAPCHKFDVKQIGAOQLLPPLYSLVFIQFVGN 60  
|||||

Db 1 MLSTSRSRFRIRNTNESGEEVTFDFDYCAPCHKFDVKQIGAOQLLPPLYSLVFIFGFGVGN 60  
Qy 61 MLVVLILINCKKLCGLTDIYLLNLAIISDLFLITPLWAHSAANEWFVGNAMCKLFTGLY 120  
Db 61 MLVVLILINCKKLCGLTDIYLLNLAIISDLFLITPLWAHSAANEWFVGNAMCKLFTGLY 120  
Qy 121 HIGVEGGFFIILLTIDRYLAIVHAFAKARTVTFGVVTSVITWLVAVFASVPGIIFTK 180  
Db 121 HIGVEGGFFIILLTIDRYLAIVHAFAKARTVTFGVVTSVITWLVAVFASVPGIIFTK 180  
Qy 181 COKEDSVVCGPYFPRGWNNEHTIMRNILGLVPLLMVICYSGILKTLRLCRNEKKRRH 240  
Db 181 COKEDSVVCGPYFPRGWNNEHTIMRNILGLVPLLMVICYSGILKTLRLCRNEKKRRH 240  
Qy 241 AVRVTFTIMVYFLEWTPYNYVILLNTQOEFGLSNCESTSDQATQVTEGLMTHCCI 300  
Db 241 AVRVTFTIMVYFLEWTPYNYVILLNTQOEFGLSNCESTSDQATQVTEGLMTHCCI 300  
Qy 301 NPILYAFVGEKPRSLFHIALGRIAPLOKPVCGGPGVPRGKNNKVVTTQGLDGRGKSKI 360  
Db 301 NPILYAFVGEKPRSLFHIALGRIAPLOKPVCGGPGVPRGKNNKVVTTQGLDGRGKSKI 360  
Qy 361 GRAPEASLQDKEGA 374  
Db 361 GRAPEASLQDKEGA 374

RESULT 4  
US-08-466-343D-9  
; Sequence 9, Application US/08466343D  
; Patent No. 6025154  
; GENERAL INFORMATION:  
; APPLICANT: LI, Yi  
; TITLE OF INVENTION: POLYNUCLEOTIDES ENCODING HUMAN G-PROTEIN  
; NUMBER OF SEQUENCES: 9  
; CORRESPONDENCE ADDRESS:  
; ADDRESSEE: STERNE, KESSLER, GOLDSTEIN & FOX P.L.L.C.  
; STREET: 1100 NEW YORK AVE., NW, SUITE 600  
; CITY: WASHINGTON  
; STATE: DC  
; COUNTRY: USA  
; ZIP: 20005  
; COMPUTER READABLE FORM:  
; MEDIUM TYPE: Floppy disk  
; COMPUTER: IBM PC compatible  
; OPERATING SYSTEM: PC-DOS/MS-DOS  
; SOFTWARE: Patent In Release #1.0, Version #1.30  
; CURRENT APPLICATION NUMBER: US/08/466,343D  
; FILING DATE: 06-JUN-1995  
; CLASSIFICATION: 435  
; ATTORNEY/AGENT INFORMATION:  
; NAME: STEFFE, ERIC K.  
; REGISTRATION NUMBER: 36,688  
; REFERENCE/DOCKET NUMBER: 1488.1150000/EKS/KLM  
; TELECOMMUNICATION INFORMATION:  
; TELEPHONE: (202) 371-2600  
; TELEFAX: (202) 371-2540  
; INFORMATION FOR SEQ ID NO: 9:  
; SEQUENCE CHARACTERISTICS:  
; LENGTH: 344 amino acids  
; TYPE: amino acid  
; STRANDEDNESS: single  
; TOPOLOGY: linear  
; MOLECULE TYPE: protein  
US-08-466-343D-9

Query Match 92.5%; Score 1823; DB 3; Length 344;  
Best Local Similarity 100.0%; Pred. No. 9, 6e-139;  
Matches 344; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 18 EVVTFDFDYCAPCHKFDVKQIGAOQLLPPLYSLVFIFGFGVGNMLVVLINCKKLCGLT 77

Db 1 EVVTFDFDYCAPCHKFDVKQIGAOQLLPPLYSLVFIFGFGVGNMLVVLINCKKLCGLT 60  
Qy 78 DYLLNLAIISDLFLITPLWAHSAANEWFVGNAMCKLFTGLYHIGYFGGIFPIILLTID 137  
Db 61 DYLLNLAIISDLFLITPLWAHSAANEWFVGNAMCKLFTGLYHIGYFGGIFPIILLTID 120  
Qy 138 RYLAIVHAFAKARTVTFGVVTSVITWLVAVFASVPGIIFTKCOKEDSVVYJCGPYFPRG 197  
Db 121 RYLAIVHAFAKARTVTFGVVTSVITWLVAVFASVPGIIFTKCOKEDSVVYJCGPYFPRG 180  
Qy 198 WNNFHTIMRNILGLVPLLMVICYSGILKTLRLCRNEKKRRHRAVRVIFTIMVYFLEW 257  
Db 181 WNNFHTIMRNILGLVPLLMVICYSGILKTLRLCRNEKKRRHRAVRVIFTIMVYFLEW 240  
Qy 258 PYNIVILLNTQOEFGLSNCESTSDQATQVTEGLMTHCCIINPIIYAFVGEKPRSLFH 317  
Db 241 PYNIVILLNTQOEFGLSNCESTSDQATQVTEGLMTHCCIINPIIYAFVGEKPRSLFH 300  
Qy 318 IALGRIAPLOKPVCGGPGVPRGKNNKVVTTQGLDGRGKSKI 361  
Db 301 IALGRIAPLOKPVCGGPGVPRGKNNKVVTTQGLDGRGKSKI 344

RESULT 5  
US-08-450-393A-4  
; Sequence 4, Application US/08450393A  
; Patent No. 5707815  
; GENERAL INFORMATION:  
; APPLICANT: Charo, Israel  
; APPLICANT: Coughlin, Shaun  
; TITLE OF INVENTION: MAMMALIAN MONOCYTE CHEMOATTRACTANT  
; NUMBER OF SEQUENCES: 14  
; CORRESPONDENCE ADDRESS:  
; ADDRESSEE: Cooley Godward Castro Huddleson & Tatum  
; STREET: 5 Palo Alto Square  
; CITY: Palo Alto  
; STATE: California  
; COUNTRY: USA  
; ZIP: 94306-2155  
; COMPUTER READABLE FORM:  
; MEDIUM TYPE: Floppy disk  
; COMPUTER: IBM PC compatible  
; OPERATING SYSTEM: PC-DOS/MS-DOS  
; SOFTWARE: Patent In Release #1.0, Version #1.25  
; CURRENT APPLICATION NUMBER: US/08/450,393A  
; FILING DATE: May 25, 1995  
; CLASSIFICATION: 424  
; ATTORNEY/AGENT INFORMATION:  
; NAME: Cserr, Luann  
; REGISTRATION NUMBER: 31,822  
; REFERENCE/DOCKET NUMBER: UCAL-237/02US  
; TELECOMMUNICATION INFORMATION:  
; TELEPHONE: 415-843-5165  
; TELEFAX: 415-8857-0663  
; INFORMATION FOR SEQ ID NO: 4:  
; SEQUENCE CHARACTERISTICS:  
; LENGTH: 360 amino acids  
; TYPE: amino acid  
; TOPOLOGY: linear  
; MOLECULE TYPE: protein  
US-08-450-393A-4

Query Match 83.8%; Score 1651.5; DB 1; Length 360;  
Best Local Similarity 95.5%; Pred. No. 5, 4e-125;  
Matches 319; Conservative 3; Mismatches 5; Indels 7; Gaps 3;

Qy 1 MLSTSRSRFRIRNTNESGEEVTFDFDYCAPCHKFDVKQIGAOQLLPPLYSLVFIFGFGVGN 60  
Db 1 MLSTSRSRFRIRNTNESGEEVTFDFDYCAPCHKFDVKQIGAOQLLPPLYSLVFIFGFGVGN 60

QY 61 MLVVLILNCKKLCITDIYLLNLAISDLFLITLPLWAHSAANEWVFGNAMCKLFTGLY 120  
DB 61 MLVVLILNCKKLCITDIYLLNLAISDLFLITLPLWAHSAANEWVFGNAMCKLFTGLY 120  
QY 121 HIGYFGGIFILLITDRYLAIHVAHFALKARTVTFGVVTSVITLWLVAVFASVPGIIFTK 180  
DB 121 HIGYFGGIFILLITDRYLAIHVAHFALKARTVTFGVVTSVITLWLVAVFASVPGIIFTK 180  
QY 181 COKEDSVYCGPYFPGWNNFHTIMRNILGLVPLLLIMVICYSGLIKTLRCRNEKKRHR 240  
DB 181 COKEDSVYCGPYFPGWNNFHTIMRNILGLVPLLLIMVICYSGLIKTLRCRNEKKRHR 240  
QY 241 AVRVIITMIVYFLWTPYNNIVILLNTFQEFFGLSNCESTSQLDQATQVTTGLMTHCCI 300  
DB 241 AVRVIITMIVYFLWTPYNNIVILLNTFQEFFGLSNCESTSQLDQATQVTTGLMTHCCI 300  
QY 301 NPIIYAFVGEKFR---SLF---HIALG-CRIAPL 327  
DB 301 NPIIYAFVGEKFR---SLF---HIALG-CRIAPL 327  
DB 301 NPIIYAFVGEKFR---SLF---HIALG-CRIAPL 327  
DB 301 NPIIYAFVGEKFR---SLF---HIALG-CRIAPL 327

## RESULT 6

US-08-446-669-4

; Sequence 4, Application US/08446669

; Patent No. 6132987

; GENERAL INFORMATION:

; APPLICANT: Charo, Israel

; TITLE OF INVENTION: MAMMALIAN MONOCYTE CHEMOATTRACTANT

; NUMBER OF SEQUENCES: 14

; CORRESPONDENCE ADDRESS:

; ADDRESSEE: Cooley Godward Castro Huddleson &amp; Tatum

; STREET: 5 Palo Alto Square

; CITY: Palo Alto

; STATE: California

; COUNTRY: USA

; ZIP: 94306-2155

; COMPUTER READABLE FORM:

; MEDIUM TYPE: Floppy disk

; COMPUTER: IBM PC compatible

; OPERATING SYSTEM: PC-DOS/MS-DOS

; SOFTWARE: Patent In Release #1.0, Version #1.25

; CURRENT APPLICATION DATA:

; APPLICATION NUMBER: US/08/446,669

; FILING DATE: May 25, 1995

; CLASSIFICATION: 435

; ATTORNEY/AGENT INFORMATION:

; NAME: Neeley, Richard

; REGISTRATION NUMBER: 30,092

; REFERENCE/DOCKET NUMBER: UCAL-237/0105

; TELECOMMUNICATION INFORMATION:

; TELEPHONE: 415-843-5000

; TELEFAX: 415-857-0663

; TELEX: 380816CooleyPA

; INFORMATION FOR SEQ ID NO: 4:

; SEQUENCE CHARACTERISTICS:

; LENGTH: 360 amino acids

; TYPE: amino acid

; TOPOLOGY: linear

; MOLECULE TYPE: protein

; FRAGMENT TYPE: internal

; US-08-446-669-4

Query Match 83.8%; Score 1651.5; DB 4; Length 360;  
Best Local Similarity 95.5%; Pred. No. 5.4e-125;  
Matches 319; Conservative 3; Mismatches 5; Indels 7; Gaps 3;

QY 1 MLSTSRSRFRINTNESGEVTFDFDYGAPCHKFDVKQIGAQQLLPPLYSLVFIQFVGN 60  
DB 1 MLSTSRSRFRINTNESGEVTFDFDYGAPCHKFDVKQIGAQQLLPPLYSLVFIQFVGN 60  
QY 61 MLVVLILNCKKLCITDIYLLNLAISDLFLITLPLWAHSAANEWVFGNAMCKLFTGLY 120

DB 61 MLVVLILNCKKLCITDIYLLNLAISDLFLITLPLWAHSAANEWVFGNAMCKLFTGLY 120  
QY 121 HIGYFGGIFILLITDRYLAIHVAHFALKARTVTFGVVTSVITLWLVAVFASVPGIIFTK 180  
DB 121 HIGYFGGIFILLITDRYLAIHVAHFALKARTVTFGVVTSVITLWLVAVFASVPGIIFTK 180  
QY 181 COKEDSVYCGPYFPGWNNFHTIMRNILGLVPLLLIMVICYSGLIKTLRCRNEKKRHR 240  
DB 181 COKEDSVYCGPYFPGWNNFHTIMRNILGLVPLLLIMVICYSGLIKTLRCRNEKKRHR 240  
QY 241 AVRVIITMIVYFLWTPYNNIVILLNTFQEFFGLSNCESTSQLDQATQVTTGLMTHCCI 300  
DB 241 AVRVIITMIVYFLWTPYNNIVILLNTFQEFFGLSNCESTSQLDQATQVTTGLMTHCCI 300  
QY 301 NPIIYAFVGEKFR---SLF---HIALG-CRIAPL 327  
DB 301 NPIIYAFVGEKFR---SLF---HIALG-CRIAPL 327  
DB 301 NPIIYAFVGEKFR---SLF---HIALG-CRIAPL 327  
DB 301 NPIIYAFVGEKFR---SLF---HIALG-CRIAPL 327

## RESULT 7

US-09-045-583-50

; Sequence 50, Application US/09045583

; Patent No. 6287805

; GENERAL INFORMATION:

; APPLICANT: Graham, Gerard J. et al.

; TITLE OF INVENTION: NO 6287805el Molecules of the G Protein-Coupled

; NUMBER OF SEQUENCES: 56

; CORRESPONDENCE ADDRESS:

; ADDRESSEE: LAHIVE &amp; COCKFIELD, LLP

; STREET: 28 State Street

; CITY: Boston

; STATE: Massachusetts

; COUNTRY: USA

; ZIP: 02109

; COMPUTER READABLE FORM:

; MEDIUM TYPE: Floppy disk

; COMPUTER: IBM PC compatible

; OPERATING SYSTEM: PC-DOS/MS-DOS

; SOFTWARE: Patent In Release #1.0, Version #1.25

; CURRENT APPLICATION DATA:

; APPLICATION NUMBER: US/09/045,583

; FILING DATE: 20-MAR-98

; CLASSIFICATION: 435

; PRIOR APPLICATION DATA:

; APPLICATION NUMBER:

; FILING DATE:

; ATTORNEY/AGENT INFORMATION:

; NAME: Mandragoras, Amy E.

; REGISTRATION NUMBER: 36,207

; REFERENCE/DOCKET NUMBER: MNI-044

; TELECOMMUNICATION INFORMATION:

; TELEPHONE: (617)227-7400

; TELEFAX: (617)742-4214

; INFORMATION FOR SEQ ID NO: 50:

; SEQUENCE CHARACTERISTICS:

; LENGTH: 360 amino acids

; TYPE: amino acid

; TOPOLOGY: linear

; MOLECULE TYPE: peptide

; FRAGMENT TYPE: internal

; US-09-045-583-50

Query Match 83.8%; Score 1651.5; DB 4; Length 360;  
Best Local Similarity 95.5%; Pred. No. 5.4e-125;  
Matches 319; Conservative 3; Mismatches 5; Indels 7; Gaps 3;

QY 1 MLSTSRSRFRINTNESGEVTFDFDYGAPCHKFDVKQIGAQQLLPPLYSLVFIQFVGN 60  
DB 1 MLSTSRSRFRINTNESGEVTFDFDYGAPCHKFDVKQIGAQQLLPPLYSLVFIQFVGN 60  
QY 61 MLVVLILNCKKLCITDIYLLNLAISDLFLITLPLWAHSAANEWVFGNAMCKLFTGLY 120

Db 61 MLVVLILNCKKLCITDIYLLNLALISDLFLITLPLWAHSAANEWVFGNAMCKLFTGLY 120  
QY 121 HIGYGGIFFIILLTDRYLAIHVAHFALKARTVTFGVVTSVITWLVAVFASVPGIIFTK 180  
Db 121 HIGYGGIFFIILLTDRYLAIHVAHFALKARTVTFGVVTSVITWLVAVFASVPGIIFTK 180  
QY 181 COKEDSVVCGPYFPGWNNFHTIMRNILGLVLPILLIMVICYSGILKTLILRCRNEKKRHR 240  
Db 181 COKEDSVVCGPYFPGWNNFHTIMRNILGLVLPILLIMVICYSGILKTLILRCRNEKKRHR 240  
QY 241 AVRVIETIMVYFLEWTPYNIIVILLNTFQEFFGLSNCSTSSOLDQATQVTTGLMTHCCI 300  
Db 241 AVRVIETIMVYFLEWTPYNIIVILLNTFQEFFGLSNCSTSSOLDQATQVTTGLMTHCCI 300  
QY 301 NPIIYAFVGEKFR---SLF---HIALG-CRIAPL 327  
Db 301 NPIIYAFVGEKFRYLSVFFRKHITRCKQCPV 334

## RESULT 8

US-09-534-185-50  
; Sequence 50, Application US/09534185  
; Patent No. 6403767  
; GENERAL INFORMATION:  
; APPLICANT: Graham, Gerard J. et al.  
; TITLE OF INVENTION: No. 6403767el Molecules of the G protein-Coupled  
; Heptahelical Receptor Superfamily and Uses  
; Therefor  
; NUMBER OF SEQUENCES: 56  
; CORRESPONDENCE ADDRESS:  
; ADDRESSEE: LAHIVE & COCKFIELD, LLP  
; STREET: 28 State Street  
; CITY: Boston  
; STATE: Massachusetts  
; COUNTRY: USA  
; ZIP: 02109  
; COMPUTER READABLE FORM:  
; MEDIUM TYPE: Floppy disk  
; COMPUTER: IBM PC compatible  
; OPERATING SYSTEM: PC-DOS/MS-DOS  
; SOFTWARE: Patentin Release #1.0, Version #1.25  
; CURRENT APPLICATION DATA:  
; APPLICATION NUMBER: US/09/534,185  
; FILING DATE: 24-Mar-2000  
; CLASSIFICATION: <Unknown>  
; PRIOR APPLICATION DATA:  
; APPLICATION NUMBER: 09/045,583  
; FILING DATE: <Unknown>  
; ATTORNEY/AGENT INFORMATION:  
; NAME: Mandragoras, Amy E.  
; REGISTRATION NUMBER: 36,207  
; REFERENCE/DOCKET NUMBER: MNI-044  
; TELECOMMUNICATION INFORMATION:  
; TELEPHONE: (617)227-7400  
; TELEFAX: (617)742-4214  
; INFORMATION FOR SEQ ID NO: 50:  
; SEQUENCE CHARACTERISTICS:  
; LENGTH: 360 amino acids  
; TYPE: amino acid  
; TOPOLOGY: linear  
; MOLECULE TYPE: peptide  
; FRAGMENT TYPE: internal  
; SEQUENCE DESCRIPTION: SEQ ID NO: 50:  
US-09-534-185-50

Query Match 83.8%; Score 1651.5; DB 4; Length 360;  
Best Local Similarity 95.5%; Pred. No. 5.4e-125;  
Matches 319; Conservative 3; Mismatches 5; Indels 7; Gaps 3;  
QY 1 MLSTSRSRFRNTNESGEEVTFDFDYDGAPCHKFDVKQIGAQLLPPLYSLVIFGFGVN 60  
Db 1 MLSTSRSRFRNTNESGEEVTFDFDYDGAPCHKFDVKQIGAQLLPPLYSLVIFGFGVN 60

QY 61 MLVVLILNCKKLCITDIYLLNLALISDLFLITLPLWAHSAANEWVFGNAMCKLFTGLY 120  
Db 61 MLVVLILNCKKLCITDIYLLNLALISDLFLITLPLWAHSAANEWVFGNAMCKLFTGLY 120  
QY 121 HIGYGGIFFIILLTDRYLAIHVAHFALKARTVTFGVVTSVITWLVAVFASVPGIIFTK 180  
Db 121 HIGYGGIFFIILLTDRYLAIHVAHFALKARTVTFGVVTSVITWLVAVFASVPGIIFTK 180  
QY 181 COKEDSVVCGPYFPGWNNFHTIMRNILGLVLPILLIMVICYSGILKTLILRCRNEKKRHR 240  
Db 181 COKEDSVVCGPYFPGWNNFHTIMRNILGLVLPILLIMVICYSGILKTLILRCRNEKKRHR 240  
QY 241 AVRVIETIMVYFLEWTPYNIIVILLNTFQEFFGLSNCSTSSOLDQATQVTTGLMTHCCI 300  
Db 241 AVRVIETIMVYFLEWTPYNIIVILLNTFQEFFGLSNCSTSSOLDQATQVTTGLMTHCCI 300  
QY 301 NPIIYAFVGEKFR---SLF---HIALG-CRIAPL 327  
Db 301 NPIIYAFVGEKFRYLSVFFRKHITRCKQCPV 334

## RESULT 9

PCT-US95-00476-4  
; Sequence 4, Application PC/TUS9500476  
; GENERAL INFORMATION:  
; APPLICANT: The Regents of the University of California  
; TITLE OF INVENTION: MAMMALIAN MONOCYTE CHEMOATTRACTANT  
; TITLE OF INVENTION: PROTEIN RECEPTORS  
; NUMBER OF SEQUENCES: 14  
; CORRESPONDENCE ADDRESS:  
; ADDRESSEE: Robbins, Berlin & Carson  
; STREET: 201 N. Figueroa Street, 5th Floor  
; CITY: Los Angeles  
; STATE: California  
; COUNTRY: USA  
; ZIP: 90012-2628  
; COMPUTER READABLE FORM:  
; MEDIUM TYPE: Floppy disk  
; COMPUTER: IBM PC compatible  
; OPERATING SYSTEM: PC-DOS/MS-DOS  
; SOFTWARE: Patentin Release #1.0, Version #1.25  
; CURRENT APPLICATION DATA:  
; APPLICATION NUMBER: PCT/US95/00476  
; FILING DATE:  
; CLASSIFICATION:  
; ATTORNEY/AGENT INFORMATION:  
; NAME: Berliner, Robert  
; REGISTRATION NUMBER: 20,121  
; REFERENCE/DOCKET NUMBER: 5555-291  
; TELECOMMUNICATION INFORMATION:  
; TELEPHONE: 310-977-1001  
; TELEFAX: 310-977-1003  
; TELEX:  
; INFORMATION FOR SEQ ID NO: 4:  
; SEQUENCE CHARACTERISTICS:  
; LENGTH: 360 amino acids  
; TYPE: amino acid  
; TOPOLOGY: linear  
; MOLECULE TYPE: protein  
; PCT-US95-00476-4

Query Match 83.8%; Score 1651.5; DB 5; Length 360;  
Best Local Similarity 95.5%; Pred. No. 5.4e-125;  
Matches 319; Conservative 3; Mismatches 5; Indels 7; Gaps 3;

QY 1 MLSTSRSRFRNTNESGEEVTFDFDYDGAPCHKFDVKQIGAQLLPPLYSLVIFGFGVN 60  
Db 1 MLSTSRSRFRNTNESGEEVTFDFDYDGAPCHKFDVKQIGAQLLPPLYSLVIFGFGVN 60  
QY 61 MLVVLILNCKKLCITDIYLLNLALISDLFLITLPLWAHSAANEWVFGNAMCKLFTGLY 120  
Db 61 MLVVLILNCKKLCITDIYLLNLALISDLFLITLPLWAHSAANEWVFGNAMCKLFTGLY 120

QY 121 HIGYGGIFFIILLTIDRYLAIVHAFKARTVTFGVVTSVITWLAVFASVPGIIFTK 180  
Db 121 HIGYGGIFFIILLTIDRYLAIVHAFKARTVTFGVVTSVITWLAVFASVPGIIFTK 180  
QY 181 CQEDSVVCGPYFPRGWNPFHTIMRNILGLVPLIMVICYSGILKTLRLCRNEKKRHR 240  
Db 181 CQEDSVVCGPYFPRGWNPFHTIMRNILGLVPLIMVICYSGILKTLRLCRNEKKRHR 240  
QY 241 AVRIFIMIVYFLFWPTYNIVILLNTFQEFFGLSNCESTSQLDQATQVETLGMTHCCI 300  
Db 241 AVRIFIMIVYFLFWPTYNIVILLNTFQEFFGLSNCESTSQLDQATQVETLGMTHCCI 300  
QY 301 NPIIYAFVGEKFR---SLF---HIALG-CRIAPL 327  
Db 301 NPIIYAFVGEKFRYLSVFFRKHHTKRFCKQCPV 334

RESULT 10  
US-08-833-752-7  
Sequence 7, Application US/08833752  
Patent No. 6448375  
GENERAL INFORMATION:  
APPLICANT: SAMSON, MICHEL  
APPLICANT: PARMENTIER, MARC  
APPLICANT: VASSART, GILBERT  
APPLICANT: LIBERT, FREDERICK  
TITLE OF INVENTION: ACTIVE AND INACTIVE CC-CHEMOKINES RECEPTOR  
NUMBER OF SEQUENCES: 17  
CORRESPONDENCE ADDRESS:  
ADDRESSEE: Knobbe, Martens, Olson & Bear  
STREET: 620 Newport Center Drive 16th Floor  
CITY: Newport Beach  
STATE: CA  
COUNTRY: U.S.A.  
ZIP: 92660  
COMPUTER READABLE FORM:  
MEDIUM TYPE: Floppy disk  
COMPUTER: IBM PC compatible  
OPERATING SYSTEM: PC-DOS/MS-DOS  
SOFTWARE: PatentIn Release #1.0, Version #1.25 (EPO)  
CURRENT APPLICATION DATA:  
APPLICATION NUMBER: US/08/833,752  
FILING DATE: 9-APR-1997  
CLASSIFICATION: 536  
ATTORNEY/AGENT INFORMATION:  
NAME: Altman, Daniel E.  
REGISTRATION NUMBER: 34,115  
REFERENCE/DOCKET NUMBER:  
INFORMATION FOR SEQ ID NO: 7:  
SEQUENCE CHARACTERISTICS:  
LENGTH: 360 amino acids  
TYPE: amino acid  
STRANDEDNESS: single  
TOPOLOGY: linear  
MOLECULE TYPE: No. 6448375e  
US-08-833-752-7

Query Match 83.5%; Score 1645.5; DB 4; Length 360;  
Best Local Similarity 94.9%; Pred. No. 1.6e-124;  
Matches 317; Conservative 4; Mismatches 6; Indels 7; Gaps 3;  
QY 1 MLSTSRFRIRNTNSESGETVTFDDYDYGAPCHKFDVKQIGALLPPLYSLVFIFGVGN 60  
Db 1 MLSTSRFRIRNTNSESGETVTFDDYDYGAPCHKFDVKQIGALLPPLYSLVFIFGVGN 60  
QY 61 MLVVLILNCKKLKCLTDIYLLNLAIISDLFLITPLWAHSAANEVFGNAMCKLFTGLY 120  
Db 61 MLVVLILNCKKLKCLTDIYLLNLAIISDLFLITPLWAHSAANEVFGNAMCKLFTGLY 120  
QY 121 HIGYGGIFFIILLTIDRYLAIVHAFKARTVTFGVVTSVITWLAVFASVPGIIFTK 180  
Db 121 HIGYGGIFFIILLTIDRYLAIVHAFKARTVTFGVVTSVITWLAVFASVPGIIFTK 180

QY 181 CQEDSVVCGPYFPRGWNPFHTIMRNILGLVPLIMVICYSGILKTLRLCRNEKKRHR 240  
Db 181 CQEDSVVCGPYFPRGWNPFHTIMRNILGLVPLIMVICYSGILKTLRLCRNEKKRHR 240  
QY 241 AVRIFIMIVYFLFWPTYNIVILLNTFQEFFGLSNCESTSQLDQATQVETLGMTHCCI 300  
Db 241 AVRIFIMIVYFLFWPTYNIVILLNTFQEFFGLSNCESTSQLDQATQVETLGMTHCCI 300  
QY 301 NPIIYAFVGEKFR---SLF---HIALG-CRIAPL 327  
Db 301 NPIIYAFVGEKFRYLSVFFRKHHTKRFCKQCPV 334

RESULT 11  
US-09-045-583-51  
Sequence 51, Application US/09045583  
Patent No. 6287805  
GENERAL INFORMATION:  
APPLICANT: Graham, Gerard J. et al.  
TITLE OF INVENTION: No. 6287805el Molecules of the G Protein-Coupled  
NUMBER OF SEQUENCES: 56  
CORRESPONDENCE ADDRESS:  
ADDRESSEE: LAHIVE & COCKFIELD, LLP  
STREET: 28 State Street  
CITY: Boston  
STATE: Massachusetts  
COUNTRY: USA  
ZIP: 02109  
COMPUTER READABLE FORM:  
MEDIUM TYPE: Floppy disk  
COMPUTER: IBM PC compatible  
OPERATING SYSTEM: PC-DOS/MS-DOS  
SOFTWARE: PatentIn Release #1.0, Version #1.25  
CURRENT APPLICATION DATA:  
APPLICATION NUMBER: US/09/045,583  
FILING DATE: 20-MAR-98  
CLASSIFICATION: 435  
PRIOR APPLICATION DATA:  
APPLICATION NUMBER:  
FILING DATE:  
ATTORNEY/AGENT INFORMATION:  
NAME: Mandragouras, Amy E.  
REGISTRATION NUMBER: 36,207  
REFERENCE/DOCKET NUMBER: MNI-044  
TELECOMMUNICATION INFORMATION:  
TELEPHONE: (617)227-7400  
TELEFAX: (617)742-4214  
INFORMATION FOR SEQ ID NO: 51:  
SEQUENCE CHARACTERISTICS:  
LENGTH: 360 amino acids  
TYPE: amino acid  
TOPOLOGY: linear  
MOLECULE TYPE: peptide  
FRAGMENT TYPE: internal  
US-09-045-583-51

Query Match 82.0%; Score 1614.5; DB 4; Length 360;  
Best Local Similarity 96.6%; Pred. No. 4.9e-122;  
Matches 308; Conservative 4; Mismatches 4; Indels 3; Gaps 1;  
QY 1 MLSTSRFRIRNTNSESGETVTFDDYDYGAPCHKFDVKQIGALLPPLYSLVFIFGVGN 60  
Db 1 MLSTSRFRIRNTNSESGETVTFDDYDYGAPCHKFDVKQIGALLPPLYSLVFIFGVGN 60  
QY 61 MLVVLILNCKKLKCLTDIYLLNLAIISDLFLITPLWAHSAANEVFGNAMCKLFTGLY 120  
Db 61 MLVVLILNCKKLKCLTDIYLLNLAIISDLFLITPLWAHSAANEVFGNAMCKLFTGLY 120  
QY 121 HIGYGGIFFIILLTIDRYLAIVHAFKARTVTFGVVTSVITWLAVFASVPGIIFTK 180  
Db 121 HIGYGGIFFIILLTIDRYLAIVHAFKARTVTFGVVTSVITWLAVFASVPGIIFTK 180

QY 181 COKEDSVYVCGPYFPGWNNFHTIMRNILGLVPLLLIMVICYSGILKTLRLCRNEKKRHR 240  
DB 181 COKEDSVYVCGPYFPGWNNFHTIMRNILGLVPLLLIMVICYSGILKTLRLCRNEKKRHR 240  
QY 241 AVRLFTIMVYFLEWTPYINIVILLNTFOEFFGLSNCESTSLDQATQVTEILGTHCCI 300  
DB 241 AVRLFTIMVYFLEWTPYINIVILLNTFOEFFGLSNCESTSLDQATQVTEILGTHCCI 300  
QY 301 NPIIYAFVGEKFR---SLF 316  
DB 301 NPIIYAFVGEKFRYLSMF 319

RESULT 12  
US-09-534-185-51  
; Sequence 51, Application US/09534185  
; Patent No. 6403767  
; GENERAL INFORMATION:  
; APPLICANT: Graham, Gerard J. et al.  
; TITLE OF INVENTION: No. 6403767el Molecules of the G protein-Coupled  
; Heptahelical Receptor Superfamily and Uses  
; Therefor  
; NUMBER OF SEQUENCES: 56  
; CORRESPONDENCE ADDRESS:  
; ADDRESSEE: LAHIVE & COCKFIELD, LLP  
; STREET: 28 State Street  
; CITY: Boston  
; STATE: Massachusetts  
; COUNTRY: USA  
; ZIP: 02109

COMPUTER READABLE FORM:  
; MEDIUM TYPE: Floppy disk  
; COMPUTER: IBM PC compatible  
; OPERATING SYSTEM: PC-DOS/MS-DOS  
; SOFTWARE: PatentIn Release #1.0, Version #1.25  
; CURRENT APPLICATION DATA:  
; FILING DATE: 24-Mar-2000  
; CLASSIFICATION: US/09/534,185  
; PRIOR APPLICATION DATA:  
; APPLICATION NUMBER: 09/045,583  
; FILING DATE: <Unknown>  
; ATTORNEY/AGENT INFORMATION:  
; NAME: Mandragouras, Amy E.  
; REGISTRATION NUMBER: 36,207  
; REFERENCE/DOCKET NUMBER: MNI-044  
; TELECOMMUNICATION INFORMATION:  
; TELEPHONE: (617)227-7400  
; TELEFAX: (617)742-4214  
; INFORMATION FOR SEQ ID NO: 51:  
; SEQUENCE CHARACTERISTICS:  
; LENGTH: 360 amino acids  
; TYPE: amino acid  
; TOPOLOGY: linear  
; MOLECULE TYPE: peptide  
; FRAGMENT TYPE: internal  
; SEQUENCE DESCRIPTION: SEQ ID NO: 51:  
US-09-534-185-51

Query Match 82.0%; Score 1614.5; DB 4; Length 360;  
Best Local Similarity 96.6%; Pred. No. 4.9e-122;  
Matches 308; Conservative 4; Mismatches 4; Indels 3; Gaps 1;

QY 1 MLSTSRFRIRNTNESGEVTTFFDYDYGAPCHKFDVKQIGAQLLPPLYSILVFTEGVGN 60  
DB 1 MLSTSRFRIRNTNESGEVTTFFDYDYGAPCHKFDVKQIGAQLLPPLYSILVFTEGVGN 60  
QY 61 MLVVLILINCKKLSLDIYLLNLAISDLLFLITPLWHAASANEWVFGNAMCKLFTGLY 120  
DB 61 MLVVLILINCKKLSLDIYLLNLAISDLLFLITPLWHAASANEWVFGNAMCKLFTGLY 120  
QY 121 HIGYFGGIFFTLLIDRYLAIVHAVFALKARTVTFGVTSVITWLVAVFASVPGIIFTK 180  
DB 121 HIGYFGGIFFTLLIDRYLAIVHAVFALKARTVTFGVTSVITWLVAVFASVPGIIFTK 180

DB 121 HIGYFGGIFFTLLIDRYLAIVHAVFALKARTVTFGVTSVITWLVAVFASVPGIIFTK 180  
QY 181 COKEDSVYVCGPYFPGWNNFHTIMRNILGLVPLLLIMVICYSGILKTLRLCRNEKKRHR 240  
DB 181 COKEDSVYVCGPYFPGWNNFHTIMRNILGLVPLLLIMVICYSGILKTLRLCRNEKKRHR 240  
QY 241 AVRLFTIMVYFLEWTPYINIVILLNTFOEFFGLSNCESTSLDQATQVTEILGTHCCI 300  
DB 241 AVRLFTIMVYFLEWTPYINIVILLNTFOEFFGLSNCESTSLDQATQVTEILGTHCCI 300  
QY 301 NPIIYAFVGEKFR---SLF 316  
DB 301 NPIIYAFVGEKFRYLSMF 319

RESULT 13  
US-08-461-244-3  
; Sequence 3, Application US/08461244  
; Patent No. 5776729  
; GENERAL INFORMATION:  
; APPLICANT: Soppet, Daniel R.  
; APPLICANT: Yi, Li  
; APPLICANT: Ruben, Steven M.  
; APPLICANT: Rosen, Craig A.  
; TITLE OF INVENTION: HUMAN G-PROTEIN RECEPTOR HGBR32  
; NUMBER OF SEQUENCES: 7  
; CORRESPONDENCE ADDRESS:  
; ADDRESSEE: CARELLA, BYRNE, BAIN, GILFILLAN, CECCHI,  
; ADDRESSEE: STUART & OLSTEIN  
; STREET: 6 Becker Farm Road  
; CITY: Roseland  
; STATE: New Jersey  
; COUNTRY: USA  
; ZIP: 07068

COMPUTER READABLE FORM:  
; MEDIUM TYPE: Floppy disk  
; COMPUTER: IBM PC compatible  
; OPERATING SYSTEM: PC-DOS/MS-DOS  
; SOFTWARE: PatentIn Release #1.0, Version #1.30  
; CURRENT APPLICATION DATA:  
; APPLICATION NUMBER: US/08/461,244  
; FILING DATE: 05-JUN-1995  
; CLASSIFICATION: 536  
; ATTORNEY/AGENT INFORMATION:  
; NAME: Feiraro, Gregory D.  
; REGISTRATION NUMBER: 36,134  
; REFERENCE/DOCKET NUMBER: 325800-445  
; TELECOMMUNICATION INFORMATION:  
; TELEPHONE: 201-994-1700  
; TELEFAX: 201-994-1744  
; INFORMATION FOR SEQ ID NO: 3:  
; SEQUENCE CHARACTERISTICS:  
; LENGTH: 347 amino acids  
; TYPE: amino acid  
; STRANDEDNESS:  
; TOPOLOGY: linear  
; MOLECULE TYPE: protein  
US-08-461-244-3

Query Match 80.7%; Score 1589.5; DB 1; Length 347;  
Best Local Similarity 95.3%; Pred. No. 4.8e-120;  
Matches 306; Conservative 3; Mismatches 5; Indels 7; Gaps 3;

QY 14 NESGEVTTFFDYDYGAPCHKFDVKQIGAQLLPPLYSILVFTEGVGNMLVLLINCKKL 73  
DB 1 NESGEVTTFFDYDYGAPCHKFDVKQIGAQLLPPLYSILVFTEGVGNMLVLLINCKKL 60  
QY 74 KCLTDIYLLNLAISDLLFLITPLWHAASANEWVFGNAMCKLFTGLYHIGYFGGIFFTIL 133  
DB 61 KCLTDIYLLNLAISDLLFLITPLWHAASANEWVFGNAMCKLFTGLYHIGYFGGIFFTIL 120  
QY 134 LTIDRYLAIVHAVFALKARTVTFGVTSVITWLVAVFASVPGIIFTKCKQEDSVVCCGPY 193  
DB 134 LTIDRYLAIVHAVFALKARTVTFGVTSVITWLVAVFASVPGIIFTKCKQEDSVVCCGPY 193

```

Db 121 LTIDRYLAIVHAFALKARTVTFGVTSVITWLVAFASVPGIIFTKCKEDSVVVCOPY 180
Qy 194 FPGWNNFHTMRNLTGLVPLLMVICYSGLKTLRCRNEKKRHRVAVRIFTIMIVYF 253
Db 181 FPGWNNFHTMRNLTGLVPLLMVICYSGLKTLRCRNEKKRHRVAVRIFTIMIVYF 240
Qy 254 LFPTPNIVILLNTFOEFFGLSNCSTSQLDQATQVTTGLMTHCCINPIIYAFVGER 313
Db 241 LFPTPNIVILLNTFOEFFGLSNCSTSQLDQATQVTTGLMTHCCINPIIYAFVGER 300
Qy 314 ---SLF---HIALG-CRIAPL 327
Db 301 RYLSVFRKHITKRECKQCPV 321

RESULT 14
US-09-517-605-5
; Sequence 5, Application US/09517605
; Patent No. 6391567
; GENERAL INFORMATION:
; APPLICANT: Littman, Dan R.
; APPLICANT: Kwon, Douglas S.
; APPLICANT: van Kooyk, Yvette
; APPLICANT: Geljenbeck, Tneo
; TITLE OF INVENTION: METHODS OF USING A FACILITATOR OF RETROVIRAL ENTRY INTO
; FILE REFERENCE: 1049-1-017
; CURRENT APPLICATION NUMBER: US/09/517,605
; CURRENT FILING DATE: 2000-03-02
; NUMBER OF SEQ ID NOS: 17
; SOFTWARE: Patentin Ver. 2.0
; SEQ ID NO 5
; LENGTH: 352
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-517-605-5

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Query Match 62.7%; Score 1236; DB 4; Length 352;
Best Local Similarity 77.4%; Pred. No. 9.8e-92;
Matches 236; Conservative 25; Mismatches 34; Indels 10; Gaps 2;

Qy 25 DYDYGAPCHKFDVKQGAQLLPPLYSLVTFVFGVGNMVLVILINCKKCLTDIYLLNL 84
Db 13 DYDTSPPCKINVKQIAARLLPPLYSLVTFVFGVGNMVLVILINCKKCLTDIYLLNL 72
Qy 85 ATSDLLFLTLPLWAHSAANEVFGNAMCKLFTGLYHIGYFGGIFFTLTIDRYLAIVH 144
Db 73 ALSDFLLTVPFWAHYAAQWDFGNTMCOLLTLGLYFIFGFSGIFFTLTIDRYLAIVH 132
Qy 145 AVFALKARTVTFGVTSVITWLVAFASVPGIIFTKCKEDSVVVCOPYF---RCWNN 200
Db 133 AVFALKARTVTFGVTSVITWLVAFASVPGIIFTKCKEDSVVVCOPYF---RCWNN 192
Qy 201 FHTMRNLTGLVPLLMVICYSGLKTLRCRNEKKRHRVAVRIFTIMIVYFLEWPPYN 260
Db 193 FQTLKIVILGLVPLLMVICYSGLKTLRCRNEKKRHRVAVRIFTIMIVYFLEWAPYN 252
Qy 261 IVILLNTFOEFFGLSNCSTSQLDQATQVTTGLMTHCCINPIIYAFVGERKFSLF---- 316
Db 253 IVILLNTFOEFFGLNCCSSNRDLQAMQVTTGLMTHCCINPIIYAFVGERKFSLF 312
Qy 317 --HIA 319
Db 313 QKHIA 317

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RESULT 15
US-08-724-984A-2
; Sequence 2, Application US/08724984A
; Patent No. 6388055
; GENERAL INFORMATION:
; APPLICANT: Derk Bergsma, Mary Brawner, and Usman Shabon
; TITLE OF INVENTION: No. 6388055el Mouse Genomic Clone of the CC-

```

```

; TITLE OF INVENTION: CKR5 Receptor
; NUMBER OF SEQUENCES: 5
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: SmithKline Beecham Corporation
; STREET: 709 Swedeland Road, P.O. Box 1539
; CITY: King of Prussia
; STATE: PA
; COUNTRY: USA
; ZIP: 19406-0939
; COMPUTER READABLE FORM:
; MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
; COMPUTER: IBM 486
; OPERATING SYSTEM: WINDOWS FOR WORKGROUPS
; SOFTWARE: MICROSOFT WORD
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/724,984A
; FILING DATE: October 3, 1996
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER:
; FILING DATE:
; ATTORNEY/AGENT INFORMATION:
; NAME: William T. Han
; REGISTRATION NUMBER: 34,344
; REFERENCE/DOCKET NUMBER: ATG50023
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 610 270 5024
; TELEFAX: 610 270 5090
; INFORMATION FOR SEQ ID NO: 2:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 354
; TYPE: Amino Acid
; TOPOLOGY: Linear
US-08-724-984A-2

Query Match 62.6%; Score 1234; DB 4; Length 354;
Best Local Similarity 74.7%; Pred. No. 1.4e-91;
Matches 230; Conservative 29; Mismatches 43; Indels 6; Gaps 2;

Qy 17 GEEVTTFFDYDYG--APCHKFDVKQGAQLLPPLYSLVTFVFGVGNMVLVILINCKK 74
Db 5 GSVPTIYDIDYGSAPCOKINVKQIAARLLPPLYSLVTFVFGVGNMVLVILINCKK 64
Qy 75 CLTDIYLLNLATSDLLFLTLPLWAHSAANEVFGNAMCKLFTGLYHIGYFGGIFFTILL 134
Db 65 SVTDIYLLNLATSDLLFLTLPLWAHSAANEVFGNAMCKLFTGLYHIGYFGGIFFTILL 124
Qy 135 TIDRYLAIVHAFALKARTVTFGVTSVITWLVAFASVPGIIFTKCKEDSVVVCOPYF 194
Db 125 TIDRYLAIVHAFALKARTVTFGVTSVITWLVAFASVPGIIFTKCKEDSVVVCOPYF 184
Qy 195 PRG-----WNNFHTIMRNLTGLVPLLMVICYSGLKTLRCRNEKKRHRVAVRIFTIMI 250
Db 185 PHTQVHFVKSFQTLKMWILSLILPLLMVICYSGLKTLRCRNEKKRHRVAVRIFTIMI 244
Qy 251 VYFLEWTPYINIVILLNTFOEFFGLSNCSTSQLDQATQVTTGLMTHCCINPIIYAFVGE 310
Db 245 VYFLEWTPYINIVILLNTFOEFFGLNCCSSNRDLQAMQATETLGMTHCCINPIIYAFVGE 304
Qy 311 KFRSLFHI 318
Db 305 KFRSLVS 312

Search completed: May 19, 2003, 16:50:01
Job time : 20.8719 secs

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